

McGREGOR et al.

U.S. National Phase of PCT/GB2003/003860

AMENDMENT TO THE SEQUENCE LISTING

Insert the attached Sequence Listing in lieu of the original.

SEQUENCE LISTING

<110> Duncan McGREGOR
 Richard ODEGRIP
 Kevin FITZGERALD
 Rosemarie HEDERER
 Bill ELDRIDGE
 Chris ULLMAN
 Philip KUHLMAN
 Davod COOMBER

<120> IN VITRO PEPTIDE EXPRESSION LIBRARY

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<141> 2005-03-04

<150> PCT/GB03/03860

<151> 2003-09-05

<150> GB 0304657.0

<151> 2003-02-28

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<151> 2003-02-27

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acg ccg aag ttc cgc gaa aaa ccg atg gaa aag gcg gtg ggc ctc acc	96
Thr Pro Lys Phe Arg Glu Lys Pro Met Glu Lys Ala Val Gly Leu Thr	
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tcc cgt ttt gat ttc gcc att cat gtg gcg cat gcc cgt tcc cgt ggt	144
Ser Arg Phe Asp Phe Ala Ile His Val Ala His Ala Arg Ser Arg Gly	
35 40 45	

ctg cgt cgg cgc atg cca ccg gtg ctg cgt cga cgg gct att gat gcg	192
Leu Arg Arg Arg Met Pro Pro Val Leu Arg Arg Arg Ala Ile Asp Ala	
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ctg ctg cag ggg ctg tgt ttc cac tat gac ccg ctg gcc aac cgc gtc	240
Leu Leu Gln Gly Leu Cys Phe His Tyr Asp Pro Leu Ala Asn Arg Val	
65 70 75 80	
cag tgt tcc atc acc aca ctg gcc att gag tgc gga ctg gcg aca gag	288
Gln Cys Ser Ile Thr Leu Ala Ile Glu Cys Gly Leu Ala Thr Glu	
85 90 95	
tcc ggt gca gga aaa ctc tcc atc acc cgt gcc acc cgg gcc ctg acg	336
Ser Gly Ala Gly Lys Leu Ser Ile Thr Arg Ala Thr Arg Ala Leu Thr	
100 105 110	
ttc ctg tca gag ctg gga ctg att acc tac cag acg gaa tat gac ccg	384
Phe Leu Ser Glu Leu Gly Leu Ile Thr Tyr Gln Thr Glu Tyr Asp Pro	
115 120 125	
ctt atc ggg tgc tac att ccg acc gac atc acg ttc aca ctg gct ctg	432
Leu Ile Gly Cys Tyr Ile Pro Thr Asp Ile Thr Phe Thr Leu Ala Leu	
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Phe Ala Ala Leu Asp Val Ser Glu Asp Ala Val Ala Ala Ala Arg Arg	
145 150 155 160	
agt cgt gtt gaa tgg gaa aac aaa cag cgc aaa aag cag ggg ctg gat	528
Ser Arg Val Glu Trp Glu Asn Lys Gln Arg Lys Lys Gln Gly Leu Asp	
165 170 175	
acc ctg ggt atg gat gag ctg ata gcg aaa gcc tgg cgt ttt gtg cgt	576
Thr Leu Gly Met Asp Glu Leu Ile Ala Lys Ala Trp Arg Phe Val Arg	
180 185 190	
gag cgt ttc cgc agt tac cag aca gag ctt cag tcc cgt gga ata aaa	624
Glu Arg Phe Arg Ser Tyr Gln Thr Glu Leu Gln Ser Arg Gly Ile Lys	
195 200 205	
cgt gcc cgt gcg cgt cgt gat gcg aac aga gaa cgt cag gat atc gtc	672
Arg Ala Arg Ala Arg Arg Asp Ala Asn Arg Glu Arg Gln Asp Ile Val	
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acc cta gtg aaa cgg cag ctg acg cgt gaa atc tcg gaa gga cgc ttc	720
Thr Leu Val Lys Arg Gln Leu Thr Arg Glu Ile Ser Glu Gly Arg Phe	
225 230 235 240	
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Thr Ala Asn Gly Glu Ala Val Lys Arg Glu Val Glu Arg Arg Val Lys	
245 250 255	
gag cgc atg att ctg tca cgt aac cgc aat tac agc cgg ctg gcc aca	816
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Ala Ser Pro
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35 40 45
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50 55 60
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65 70 75 80
Gln Cys Ser Ile Thr Thr Leu Ala Ile Glu Cys Gly Leu Ala Thr Glu
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Glu Arg Phe Arg Ser Tyr Gln Thr Glu Leu Gln Ser Arg Gly Ile Lys
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Arg Ala Arg Ala Arg Arg Asp Ala Asn Arg Glu Arg Gln Asp Ile Val
210 215 220

Thr Leu Val Lys Arg Gln Leu Thr Arg Glu Ile Ser Glu Gly Arg Phe
 225 230 235 240
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26

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Lys His

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Asn Phe

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Ala Asp

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